

(#5)

RAW SEQUENCE LISTING

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Application Serial Number: 101531,965
Source: PCT
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PATENT APPLICATION: US/10/531,965

DATE: 04/27/2005

TIME: 14:36:39

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Output Set: N:\CRF4\04272005\J531965.raw

3 <110> APPLICANT: Bayer AG, BHC
 4 Golz, Stefan
 5 Bruggemeier, Ulf
 6 Geerts, Andreas
 8 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases Associated with

Human

9 Phosphodiesterase 10A (PDE10A)

11 <130> FILE REFERENCE: Le A 36 374

C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/531,965

C--> 13 <141> CURRENT FILING DATE: 2005-04-19

13 <150> PRIOR APPLICATION NUMBER: PCT/EP2003/011882

14 <151> PRIOR FILING DATE: 2003-10-25

16 <150> PRIOR APPLICATION NUMBER: EP02024996.7

17 <151> PRIOR FILING DATE: 2002-11-08

19 <160> NUMBER OF SEQ ID NOS: 5

21 <170> SOFTWARE: PatentIn version 3.3

23 <210> SEQ ID NO: 1

24 <211> LENGTH: 4576

25 <212> TYPE: DNA

26 <213> ORGANISM: Homo sapiens

28 <400> SEQUENCE: 1

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33	gctgttccg aaggctgacc gagtgttcc tgagcccaa agctaattgtt atagatata	180
35	aagtcatcca cagagatgtt acagttgaag agatgggggt agagaagact ttgaaggaaa	240
37	agaatgtaga atgaggatag aagagaggaa atcccaacat ttaacaggtt tgacagatga	300
39	aaaagtgaag gcataatctt ctcttcaccc ccaggttata gatgaatttg tatctgaaag	360
41	tgttagtgca gagacagtag agaaatggct gaagaggaag aacaacaaat cagaagatga	420
43	atcagctcct aaggaagtca gcaggtacca agatacgaat atgcaggggag ttgtatata	480
45	actaaacagc tatataaaac aacgggttggc cacaggagga gacaaccagc tactcctcta	540
47	tgaactgagc agcatcatta aaatagccac aaaagccgat ggatttgcac tgtatccct	600
49	tggagagtgc aataatagcc tttgtatatt cacgcaccc gggataaaagg aaggaaaacc	660
51	ccgcctcattc cctgctggc ccatcactca gggcaccacc gtctctgc ttatgtggccaa	720
53	gtccaggaaa acactgctag tagaagacat cttggagat gaacgatttc caagaggtac	780
55	tggactggaa tcagggactc gtatccagtc ttgtttgc ttaccaatttgc tcactgcaat	840
57	tgttgacttg attggatttc tcgagctgtc tcggcactgg ggcacaaagaag cttctgtct	900
59	tagtcaccag gaggttgcac cagcaaatct tgcctggct tcagtagcaa tacatcaggt	960
61	gcaggtatgc agaggcattt ccaaacagac agaattgaat gacttcctac tcgacgtatc	1020
63	aaaaaacatat tttgataaca tagttgcaat agattctcta cttgaacaca taatgatata	1080
65	tgcacaaaaac ctggtaatgc ccgatcggtt tgccgttttc caggtggacc ataagaacaa	1140
67	ggagttatat tcagacctt ttgtatattgg agagggaaaag gaagggaaaac ctgtcttcaa	1200
69	gaagacccaaa gagataagat ttcaatttgc gaaaggaatt gctggccaa tagcaagaac	1260
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79	gtatcataga	attcgccact	cagagtgc	at	ttaccggta	acgatggaaa	agctgtc	ccta	1560											
81	ccatagcatt	tgtacttcag	aagagtggca	agg	tctcatg	caattc	accc	ttcc	cg	1620										
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95	cacttccacc	atggagc	gac	accactt	c	ccagact	gtg	tccat	cctt	agttggagg	2040									
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111	cacgg	gc	c	tt	ct	gaa	ag	cat	gc	agg	ga	ta	ct	c	at	ct	ca	gt	2520	
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173 cactatgagg taaaatgtaa taacaaaaga gagagaagta ccactgtggc tagatatata 4380
 175 cacacacata tatatatgga tggatgtaat atatgttagca cacacacata gatgtatata 4440
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 196 20 25 30
 199 Phe Val Ser Glu Ser Val Ser Ala Glu Thr Val Glu Lys Trp Leu Lys
 200 35 40 45
 203 Arg Lys Asn Asn Lys Ser Glu Asp Glu Ser Ala Pro Lys Glu Val Ser
 204 50 55 60
 207 Arg Tyr Gln Asp Thr Asn Met Gln Gly Val Val Tyr Glu Leu Asn Ser
 208 65 70 75 80
 211 Tyr Ile Glu Gln Arg Leu Asp Thr Gly Gly Asp Asn Gln Leu Leu Leu
 212 85 90 95
 215 Tyr Glu Leu Ser Ser Ile Ile Lys Ile Ala Thr Lys Ala Asp Gly Phe
 216 100 105 110
 219 Ala Leu Tyr Phe Leu Gly Glu Cys Asn Asn Ser Leu Cys Ile Phe Thr
 220 115 120 125
 223 Pro Pro Gly Ile Lys Glu Gly Lys Pro Arg Leu Ile Pro Ala Gly Pro
 224 130 135 140
 227 Ile Thr Gln Gly Thr Thr Val Ser Ala Tyr Val Ala Lys Ser Arg Lys
 228 145 150 155 160
 231 Thr Leu Leu Val Glu Asp Ile Leu Gly Asp Glu Arg Phe Pro Arg Gly
 232 165 170 175
 235 Thr Gly Leu Glu Ser Gly Thr Arg Ile Gln Ser Val Leu Cys Leu Pro
 236 180 185 190
 239 Ile Val Thr Ala Ile Gly Asp Leu Ile Gly Ile Leu Glu Leu Tyr Arg
 240 195 200 205
 243 His Trp Gly Lys Glu Ala Phe Cys Leu Ser His Gln Glu Val Ala Thr
 244 210 215 220
 247 Ala Asn Leu Ala Trp Ala Ser Val Ala Ile His Gln Val Gln Val Cys
 248 225 230 235 240
 251 Arg Gly Leu Ala Lys Gln Thr Glu Leu Asn Asp Phe Leu Leu Asp Val
 252 245 250 255
 255 Ser Lys Thr Tyr Phe Asp Asn Ile Val Ala Ile Asp Ser Leu Leu Glu
 256 260 265 270
 259 His Ile Met Ile Tyr Ala Lys Asn Leu Val Asn Ala Asp Arg Cys Ala
 260 275 280 285
 263 Leu Phe Gln Val Asp His Lys Asn Lys Glu Leu Tyr Ser Asp Leu Phe
 264 290 295 300
 267 Asp Ile Gly Glu Lys Glu Gly Lys Pro Val Phe Lys Lys Thr Lys

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275	Thr	Gly	Glu	Val	Leu	Asn	Ile	Pro	Asp	Ala	Tyr	Ala	Asp	Pro	Arg	Phe
276							340			345						350
279	Asn	Arg	Glu	Val	Asp	Leu	Tyr	Thr	Gly	Tyr	Thr	Thr	Arg	Asn	Ile	Leu
280							355			360						365
283	Cys	Met	Pro	Ile	Val	Ser	Arg	Gly	Ser	Val	Ile	Gly	Val	Val	Gln	Met
284							370			375						380
287	Val	Asn	Lys	Ile	Ser	Gly	Ser	Ala	Phe	Ser	Lys	Thr	Asp	Glu	Asn	Asn
288							385			390			395			400
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295	Met	Tyr	His	Arg	Ile	Arg	His	Ser	Glu	Cys	Ile	Tyr	Arg	Val	Thr	Met
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303	Leu	Met	Gln	Phe	Thr	Leu	Pro	Val	Arg	Leu	Cys	Lys	Glu	Ile	Glu	Leu
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311	Val	Tyr	Met	Val	His	Arg	Ser	Cys	Gly	Thr	Ser	Cys	Phe	Glu	Leu	Glu
312								485			490					495
315	Lys	Leu	Cys	Arg	Phe	Ile	Met	Ser	Val	Lys	Lys	Asn	Tyr	Arg	Arg	Val
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319	Pro	Tyr	His	Asn	Trp	Lys	His	Ala	Val	Thr	Val	Ala	His	Cys	Met	Tyr
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323	Ala	Ile	Leu	Gln	Asn	Asn	His	Thr	Leu	Phe	Thr	Asp	Leu	Glu	Arg	Lys
324								530			535					540
327	Gly	Leu	Leu	Ile	Ala	Cys	Leu	Cys	His	Asp	Leu	Asp	His	Arg	Gly	Phe
328								545			550			555		560
331	Ser	Asn	Ser	Tyr	Leu	Gln	Lys	Phe	Asp	His	Pro	Leu	Ala	Ala	Leu	Tyr
332								565			570					575
335	Ser	Thr	Ser	Thr	Met	Glu	Gln	His	His	Phe	Ser	Gln	Thr	Val	Ser	Ile
336								580			585					590
339	Leu	Gln	Leu	Glu	Gly	His	Asn	Ile	Phe	Ser	Thr	Leu	Ser	Ser	Ser	Glu
340								595			600					605
343	Tyr	Glu	Gln	Val	Leu	Glu	Ile	Ile	Arg	Lys	Ala	Ile	Ile	Ala	Thr	Asp
344								610			615					620
347	Leu	Ala	Leu	Tyr	Phe	Gly	Asn	Arg	Lys	Gln	Leu	Glu	Glu	Met	Tyr	Gln
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351	Thr	Gly	Ser	Leu	Asn	Leu	Asn	Asn	Gln	Ser	His	Arg	Asp	Arg	Val	Ile
352								645			650					655
355	Gly	Leu	Met	Met	Thr	Ala	Cys	Asp	Leu	Cys	Ser	Val	Thr	Lys	Leu	Trp
356								660			665					670
359	Pro	Val	Thr	Lys	Leu	Thr	Ala	Asn	Asp	Ile	Tyr	Ala	Glu	Phe	Trp	Ala
360								675			680					685
363	Glu	Gly	Asp	Glu	Met	Lys	Lys	Leu	Gly	Ile	Gln	Pro	Ile	Pro	Met	Met
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368 705 710 715 720
371 Asn Ala Val Ala Ile Pro Cys Tyr Thr Thr Leu Thr Gln Ile Leu Pro
372 725 730 735
375 Pro Thr Glu Pro Leu Leu Lys Ala Cys Arg Asp Asn Leu Ser Gln Trp
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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date